

Figure 1A

M Q R L G A T L L C	10
GGCACGAGGGGGCGCGCTGCCGGCGCAGAGCGGAG ATG CAG CGG CTT GGG GCC ACC CTG CTG TGC	67
L L L A A A V P T A P A P A P T A T S A	30
CTG CTG CTG GCG GCG GCG GTC CCC ACG GCC CCC GCG CCC GCT CCG ACG GCG ACC TCG GCT	127
P V K P G P A L S Y P Q E E A T L N E M	50
CCA GTC AAG CCC GGC CCG GCT CTC AGC TAC CCG CAG GAG GAG GCC ACC CTC AAT GAG ATG	187
F R E V E E L M E D T Q H K L R S A V E	70
TTC CGC GAG GTT GAG GAA CTG ATG GAG GAC ACG CAG CAC AAA TTG CGC AGC GCG GTG GAA	247
E M E A E E A A A K A S S E V N L A N L	90
GAG ATG GAG GCA GAA GAA GCT GCT AAA GCA TCA TCA GAA GTG AAC CTG GCA AAC TTA	307
P P S Y H N E T N T D T N V G N N T I H	110
CCT CCC AGC TAT CAC AAT GAG ACC AAC ACA GAC ACG AAC GTT GGA AAT AAT ACC ATC CAT	367
V H R E I H K I T N N Q T G Q M V F S E	130
GTG CAC CGA GAA ATT CAC AAG ATA ACC AAC AAC CAG ACT GGA CAA ATG GTC TTT TCA GAG	427
T V I T S V G D E E G R R S H E C I I D	150
ACA GTT ATC ACA TCT GTG GGA GAC GAA GGC AGA AGG AGC CAC GAG TGC ATC ATC GAC	487
E D C G P S M Y C Q F A S F Q Y T C Q P	170
GAG GAC TGT GGG CCC AGC ATG TAC TGC CAG TTT GCC AGC TTC CAG TAC ACC TGC CAG CCA	547
C R G Q R M L C T R D S E C C G D Q L C	190
TGC CGG GGC CAG AGG ATG CTC TGC ACC CGG GAC AGT GAG TGC TGT GGA GAC CAG CTG TGT	607
V W G H C T K M A T R G S N G T I C D N	210
GTC TGG GGT CAC TGC ACC AAA ATG GCC ACC AGG GGC AGC AAT GGG ACC ATC TGT GAC AAC	667
Q R D C Q P G L C C A F Q R G L L F P V	230
CAG AGG GAC TGC CAG CCG GGG CTG TGC TGT GCC TTC CAG AGA GGC CTG CTG TTC CCT GTG	727
C T P L P V E G E L C H D P A S R L L D	250
TGC ACA CCC CTG CCC GTG GAG GGC GAG CTT TGC CAT GAC CCC GCC AGC CGG CTT CTG GAC	787
L I T W E L E P D G A L D R C P C A S G	270
CTC ATC ACC TGG GAG CTA GAG CCT GAT GGA GCC TTG GAC CGA TGC CCT TGT GCC AGT GGC	847
L L C Q P H S H S L V Y V C K P T F V G	290
CTC CTC TGC CAG CCC CAC AGC CAC AGC CTG GTG TAT GTG TGC AAG CCG ACC TTC GTG GGG	907
S R D Q D G E I L L P R E V P D E Y E V	310
AGC CGT GAC CAA GAT GGG GAG ATC CTG CTG CCC AGA GAG GTC CCC GAT GAG TAT GAA GTT	967
G S F M E E V R Q E L E D L E R S L T E	330
GGC AGC TTC ATG GAG GAG GTG CGC CAG GAG CTG GAG GAC CTG GAG AGG AGC CTG ACT GAA	1027
E M A L L E P A A A A A A L L G R E E I	350
GAG ATG GCG CTG AAG GAG CCT GCG GCT GCC GCG GCT GCA CTG CTG GGA AGG GAA GAG ATT	1087
*	351
TAG	1090

Figure 1B

ATCTGGACCAGGCTGGGTAGATGTCAATAGAAATAGCTAATTATTCCTCANGTGTGCTTTAACGGTGGCTG 1169
ACCAGGCTTCTTCCCTACATCTTCCCAGTAAGTTCCCTCTGGCTTGACAGCATGAGGTGTTGCATTTGTCAG 1248
CTCCCCCAGGCTGTTCTCCAGGCTCACAGTCTGGTCTGGAGAGTCAGGCAGGGTAAACTGCAGGAGCAGTTGC 1327
CACCCCTGTCCAGATTATGGCTGCTTGCCTCTACCAGTGGCAGACAGCCGTTGTTCTACATGGCTTGATAATTG 1406
TTTGAGGGGAGGAGATGGAAACAATGTGGAGTCTCCCTCTGATTGGTTGGGAAATGTGGAGAAGAGTGCCCTGCTT 1485
TGCAAAACATCAACCTGGAAAAATGCAACAAATGAATTTCACGCAGTTCTTCATGGCATAGGTAAGCTGTGCC 1564
TCAGCTGTCAGATGAAATGTTCTGTTCACCCCTGCATTACATGTGTTATTCATCCAGCAGTGTGCTCAGCTCCTAC 1643
CTCTGTGCCAGGGCAGCATTTCATATCCAAGATCAATTCCCTCTCAGCACAGCCTGGGAGGGGGTATTGTTCTC 1722
CTCGTCCATCAGGGATTCAGAGGCTCAGAGACTGCAAGCTGCTGCCAAGTCACACAGCTAGTGAAGACCAGAGCAG 1801
TTTCATCTGGTTGTGACTCTAAGCTCAGTGCTCTCCACTACCCACACCAGCCTGGTGCCACAAAAGTGCTCCCC 1880
AAAAGGAAGGAGAATGGGATTTCTTTGAGGCATGCACATCTGAATTAGGTCAAACATAATTCTCACATCCCTCTA 1959
AAAGTAAACTACTGTTAGAACAGCAGTGTTCTCACAGTGTGGGCAGCCGCCTCTAATGAAGACAATGATATTGAC 2038
ACTGTCCTCTTGCACTGATTAGTAACCTTGAAAGGTATATGACTGAGCGTAGCATAAGGTTAACCTGCAGAAA 2117
CAGTACTTAGGTAATTGTAGGGCAGGATTATAATGAAATTGCAAAATCACTTAGCAGCAACTGAAGACAATTATCA 2196
ACCACGTGGAGAAAATCAAACCGAGCAGGGCTGTGAAACATGGTTGAAATATGCGACTGCGAACACTGAACCTACG 2275
CCACTCCACAAATGATGTTTCAGGTGTCATGGACTGTTGCCACCATGTATTCCAGAGTTCTAAAGTTAAAGTT 2354
GCACATGATTGTATAAGCATGCTTCTTGAGTTAAATTATGATAAACATAAGTTGCATTAGAAATCAAGCATAA 2433

ATCACTTCAACTGCTAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2479

Figure 2

GAATTCCGGCACGAGAGACGACGTGCTGAGCTGCCAGCTTAGTGGAAAGCTCTGGCTCTGGGTGGAGAGCAGCCTCGCTTTG 79
 GTGACGCACAGTGTGGACCCCTCCAGGAGCCCCGGGATTGAAGG ATG GTG GCG GCC GTC CTG CTG GGG 8
 L S W L C S P L G A L V L D F N N I R S 8
 CTG AGC TGG CTC TGC TCT CCC CTG GGA GCT CTG GTC CTG GAC TTC AAC AAC ATC AGG AGC 148
 S A D L H G A R K G S Q C L S D T D C N 28
 TCT GCT GAC CTG CAT GGG GCC CGG AAG GGC TCA CAG TGC CTG TCT GAC ACG GAC TGC AAT 208
 T R K F C L Q P R D E K P F C A T C R G 48
 ACC AGA AAG TTC TGC CTC CAG CCC CGC GAT GAG AAG CCG TTC TGT GCT ACA TGT CGT GAC AGC TGC AAT 268
 L R R R C Q R D A M C C P G T L C V N D 68
 TTG CGG AGG AGG TGC CAG CGA GAT GCC ATG TGC CCT GGG ACA CTC TGT GTG AAC GAT 328
 V C T T M E D A T P I L E R Q L D E Q D 88
 GTT TGT ACT ACG ATG GAA GAT GCA ACC CCA ATA TTA GAA AGG CAG CTT GAT GAG CAA GAT 388
 G T H A E G T T G H P V Q E N Q P K R K 108
 GGC ACA CAT GCA GAA GGA ACA ACT GGG CAC CCA GTC CAG GAA AAC CAA CCC AAA AGG AAG 448
 P S I K K S Q G R K G Q E G E S C L R T 128
 CCA AGT ATT AAG AAA TCA CAA GGC AGG AAG GGA CAA GAG GGA GAA AGT TGT CTG AGA ACT 508
 F D C G P G L C C A R H F W T K I C K P 148
 TTT GAC TGT GGC CCT GGA CTT TGC TGT GCT CGT CAT TTT TGG ACG AAA ATT TGT AAG CCA 568
 V L L E G Q V C S R R G H K D T A Q A P 168
 GTC CTT TTG GAG GGA CAG GTC TGC TCC AGA AGA GGG CAT AAA GAC ACT GCT CAA GCT CCA 628
 E I F Q R C D C G P G L L C R S Q L T S 188
 GAA ATC TTC CAG CGT TGC GAC TGT GGC CCT GGA CTA CTG TGT CGA AGC CAA TTG ACC AGC 688
 N R Q H A R L R V C Q K I E K L * 208
 AAT CGG CAG CAT GCT CGA TTA AGA GTA TGC CAA AAA ATA GAA AAG CTA TAA 748
 ATATTTCAAAATAAAGAAGAATCCACATTGCAAAAAAAAAAAAAAAA 225
 799
 848

Figure 3A

CCGGACCGCGTGGCGGCACGGTTCTGTGGGGACCCAGGCTTGCAAAGTGACGGTCATTTCTCTTCTCCCTCTT 79
 M M A L G A A G A T R V F V A M
 GAGTCCTCTGAG ATG ATG GCT CTG GGC GCA GCG GGA GCT ACC CGG GTC TTT GTC GCG ATG 16
 140
 V A A A A L G G H P L L G V S A T L N S V
 GTA GCG GCG GCT CTC GGC GGC CAC CCT CTG CTG GGA GTG AGC GCC ACC TTG AAC TCG GTT 36
 200
 L N S N A I K N L P P P L G G A A G H P
 CTC AAT TCC AAC GCT ATC AAG AAC CTG CCC CCA CCG CTG GGC GGC GCT GCG GGG CAC CCA 56
 260
 G S A V S A A P G I L Y P G G N K Y Q T
 GGC TCT GCA GTC AGC GCC GCG CCG GGA ATC CTG TAC CCG GGC GGG AAT AAG TAC CAG ACC 76
 320
 I D N Y Q P Y P C A E D E E C G T D E Y
 ATT GAC AAC TAC CAG CCG TAC CCG TGC GCA GAG GAC GAG GAG TGC GGC ACT GAT GAG TAC 96
 380
 C A S P T R G G D A G V Q I C L A C R K 116
 TGC GCT AGT CCC ACC CGC GGA GGG GAC GCA GGC GTG CAA ATC TGT CTC GCC TGC AGG AAG 440
 R R K R C M R H A M C C P G N Y C K N G 136
 CGC CGA AAA CGC TGC ATG CGT CAC GCT ATG TGC TGC CCC GGG AAT TAC TGC AAA AAT GGA 500
 I C V S S D Q N H F R G E I E E T I T E 156
 ATA TGC GTG TCT TCT GAT CAA AAT CAT TTC CGA GGA GAA ATT GAG GAA ACC ATC ACT GAA 560
 S F G N D H S T L D G Y S R R T T L S S 176
 AGC TTT GGT AAT GAT CAT AGC ACC TTG GAT GGG TAT TCC AGA AGA ACC ACC TTG TCT TCA 620
 K M Y H T K G Q E G S V C L R S S D C A 196
 AAA ATG TAT CAC ACC AAA GGA CAA GAA GGT TCT GTT TGT CTC CGG TCA TCA GAC TGT GCC 680
 S G L C C A R H F W S K I C K P V L K E 216
 TCA GGA TTG TGT TGT GCT AGA CAC TTC TGG TCC AAG ATC TGT AAA CCT GTC CTG AAA GAA 740
 G Q V C T K H R R K G S H G L E I F Q R 236
 GGT CAA GTG TGT ACC AAG CAT AGG AGA AAA GGC TCT CAT GGA CTA GAA ATA TTC CAG CGT 800
 C Y C G E G L S C R I Q K D H H Q A S N 256
 TGT TAC TGT GGA GAA GGT CTG TCT TGC CGG ATA CAG AAA GAT CAC CAT CAA GCC AGT AAT 860
 S S R L H T C Q R H *
 TCT TCT AGG CTT CAC ACT TGT CAG AGA CAC TAA 267
 893
 ACCAGCTATCCAAAATGCAGTGAACCTCTTTATATAATAGATGCTATGAAAACCTTTATGACCTCATCAACTCAAT 972
 CCTAAGGATATACAAGTCTGTGGTTCTAGTTAACATTCAATAACACCTCCAAAAACCTGGAGTCTAAGAGCTTG 1051
 TTTCTTTATGGAACTCCCCGTGATTGCAGTAAATTACTGTATTGTAATTCTCAGTGTGGCACTTACCTGTAAATGCA 1130
 ATGAAACTTTAATTATTTCTAAAGGTGCTGCAC TGCTATTTCTCTTGTATGTAATTGTACACATTGA 1209
 TTGTTATCTGACTGACAAATATTCTATATTGAACTGAAGTAAATCATTCAAGCTTATAGTTCTAAAAGCATAACCCT 1288
 TTACCCCATTNATTCTAGAGTCNAGAACGCAAGGATCTCTTGAATGACAAATGATAGGTACCTAAAATGTAACATGA 1367

Figure 3B

AAATACTAGCTTATTTCTGAAATGTACTATCTTAATGCTTAAATTATATTCCTTAGGCTGTGATAGTTTGAAA 1446
TAAAATTTAACATTTAATATCATGAAATGKTATAAGTAGACATAAAAAAAAAAAAAAGGGCGGCCGCTAGA 1525
CTAG 1529

Figure 4

E	F	G	T	R	V	G	R	Y	C	H	S	P	H	Q	G	S	S	A	C	20
GAA	TTC	GGC	ACG	AGG	GTT	GGG	AGG	TAT	TGC	CAC	AGT	CCC	CAC	CAA	GGA	TCA	TCG	GCC	TGC	60
M	V	C	R	R	K	K	K	R	C	H	R	D	G	M	C	C	P	S	T	40
ATG	GTG	TGT	CGG	AGA	AAA	AAG	AAG	CGC	TGC	CAC	CGA	GAT	GGC	ATG	TGC	TGC	CCC	AGT	ACC	120
R	C	N	N	G	I	C	I	P	V	T	E	S	I	L	T	P	H	I	P	60
CGC	TGC	AAT	AAT	GGC	ATC	TGT	ATC	CCA	GTT	ACT	GAA	AGC	ATC	TTA	ACC	CCT	CAC	ATC	CCG	180
A	L	D	G	T	R	H	R	D	R	N	H	G	H	Y	S	N	H	D	L	80
GCT	CTG	GAT	GGT	ACT	CGG	CAC	AGA	GAT	CGA	AAC	CAC	GGT	CAT	TAC	TCA	AAC	CAT	GAC	TTG	240
G	W	Q	N	L	G	R	P	H	T	K	M	S	H	I	K	G	H	E	G	100
GGA	TGG	CAG	AAT	CTA	GGG	AGA	CCA	CAC	ACT	AAG	ATG	TCA	CAT	ATA	AAA	GGG	CAT	GAA	GGA	300
D	P	C	L	R	S	S	D	C	I	E	G	F	C	C	A	R	H	F	W	120
GAC	CCC	TGC	CTA	CGA	TCA	TCA	GAC	TGC	ATT	GAA	GGG	TTT	TGC	TGT	GCT	CGT	CAT	TTC	TGG	360
T	K	I	C	K	P	V	L	H	Q	G	E	V	C	T	K	Q	R	K	K	140
ACC	AAA	ATC	TGC	AAA	CCA	GTG	CTC	CAT	CAG	GGG	GAA	GTC	TGT	ACC	AAA	CAA	CGC	AAG	AAG	420
G	S	H	G	L	E	I	F	Q	R	C	D	C	A	K	G	L	S	C	K	160
GGT	TCT	CAT	GGG	CTG	GAA	ATT	TTC	CAG	CGT	TGC	GAC	TGT	GCG	AAG	GGC	CTG	TCT	TGC	AAA	480
V	W	K	D	A	T	Y	S	S	K	A	R	L	H	V	C	Q	K	I	*	180
GTA	TGG	AAA	GAT	GCC	ACC	TAC	TCC	TCC	AAA	GCC	AGA	CTC	CAT	GTG	TGT	CAG	AAA	ATT	TGA	540
TCACCATTGAGGAACATCATCAATTGCAGACTGTGAAGTGTGTATTTAATGCATTATAGCATGGTGGAAAATAAGGTT	619																			
CAGATGCAGAAGAATGGCTAAAATAAGAACGTGATAAGAATATAGATGATCACAAAAAAAAAAAAAAGATGCGG	698																			
CCGC	702																			

Figure 5

M	1
CTCGAGGCCAAATTGGCACGAGGCCGGCTGGTCTAGCATAAAGGCGGAGCCCAGAAGAAGGGCGGGGT	ATG 77
G E A S P P A P A R R H L L L V L L L L L L	21
GGA GAA GCC TCC CCA CCT GCC CCC GCA AGG CGG CAT CTG CTG GTC CTG CTG CTG CTC CTC	137
S T L V I P S A A A P I H D A D A Q E S	41
TCT ACC CTG GTG ATC CCC TCC GCT GCA GCT CCT ATC CAT GAT GCT GAC GCC CAA GAG AGC	197
S L G L T G L Q S L L Q G F S R L F L K	61
TCC TTG GGT CTC ACA GGCG CTC CAG AGC CTA CTC CAA GGCG TTC AGC CGA CTT TTC CTG AAA	257
G N L L R G I D S L F S A P M D F R G L	81
GGT AAC CTG CTT CGG GGCG ATA GAC AGC TTA TTC TCT GCC CCC ATG GAC TTC CGG GGCG CTC	317
P G N Y H K E E N Q E H Q L G N N T L S	101
CCT GGG AAC TAC CAC AAA GAG GAG AAC CAG GAG CAC CAG CTG GGG AAC AAC ACC CTC TCC	377
S H L Q I D K M T D N K T G E V L I S E	121
AGC CAC CTC CAG ATC GAC AAG ATG ACC GAC AAC AAG ACA GGA GAG GTG CTG ATC TCC GAG	437
N V V A S I Q P A E G S F E G D L K V P	141
AAT GTG GTG GCA TCC ATT CAA CCA GCG GAG GGG AGC TTC GAG GGT GAT TTG AAG GTA CCC	497
R M E E K E A L V P I Q K A T D S F H T	161
AGG ATG GAG GAG AAG GAG GCC CTG GTA CCC ATC CAG AAG GCC ACG GAC AGC TTC CAC ACA	557
E L H P R V A F W I I K L P R R R S H Q	181
GAA CTC CAT CCC CGG GTG GCC TTC TGG ATC ATT AAG CTG CCA CGG CGG AGG TCC CAC CAG	617
D A L E G G H W L S E K R H R L Q A I R	201
GAT GCC CTG GAG GGCG CAC TGG CTC AGC GAG AAG CGA CAC CGC CTG CAG GCC ATC CGG	677
D G L R K G T H K D V L E E G T E S S S	221
GAT GGA CTC CGC AAG GGG ACC CAC AAG GAC GTC CTA GAA GAG GGG ACC GAG AGC TCC TCC	737
H S R L S P R K T H L L Y I L R P S R Q	241
CAC TCC AGG CTG TCC CCC CGA AAG ACC CAC TTA CTG TAC ATC CTC AGG CCC TCT CGG CAG	797
L *	243
CTG TAG	803
GGGTGGGGACGGGGAGCACCTGCCTGTAGCCCCATCAGACCCCTGCCCAAGCACCATATGAAATAAAGTTCTTCT	882
TACATCTAAAAAAAAAAAAAAAATGGCGGCCGC	928

Figure 6

Figure 7

The Human CRSP Family

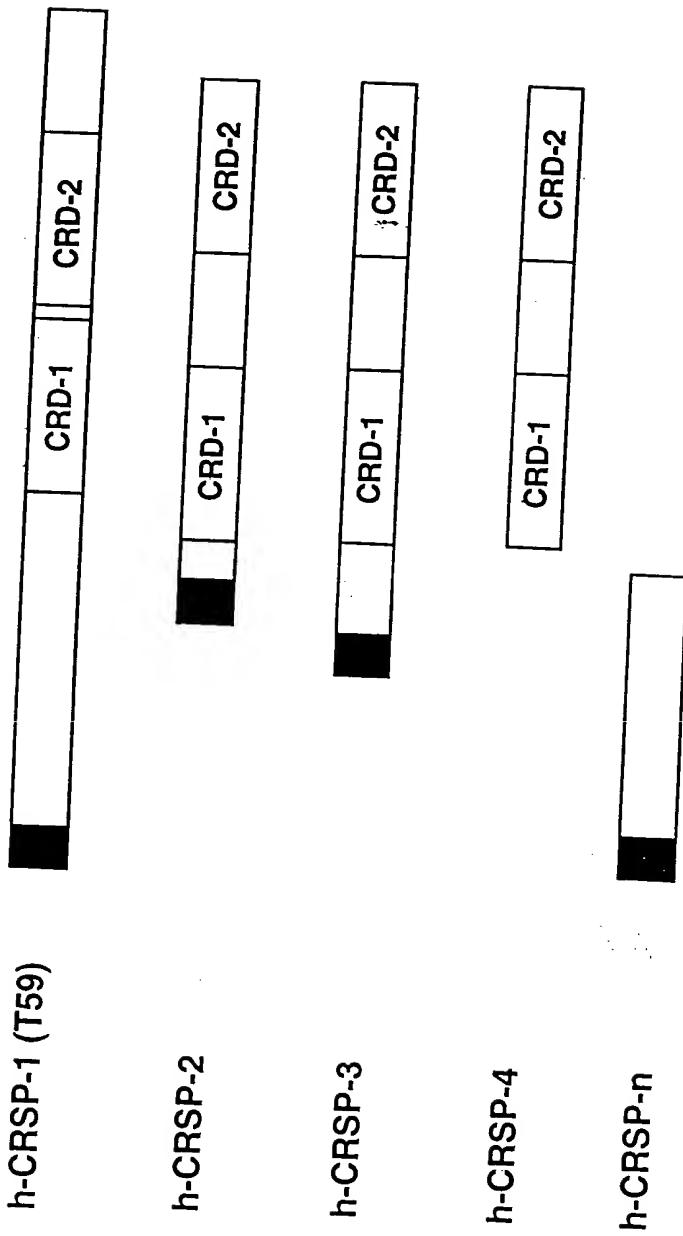


Figure 8A

FGTCGACCCACGCGTCCGCTGTGGCAGCCCAGCTACCGTGTGACCAGATCCAGCTTGAGCTCAGCTTGTTCATTC	79
M Q R L G G I L L C T L	12
GAATTGGCGGGCCAGCGCGAACAAAC ATG CAG CGG CTC GGG GGT ATT TTG CTG TGT ACA CTG	145
L A A A V P T A P A P S P T V T W T P A	32
CTG GCG GCG GCG GTC CCC ACT GCT CCT GCT CCT TCC CCG ACG GTC ACT TGG ACT CCG GCG	205
E P G P A L N Y P Q E E A T L N E M F R	52
GAG CCG GGC CCA GCT CTC AAC TAC CCT CAG GAG GAA GCT ACG CTC AAT GAG ATG TTT CGA	265
E V E E L M E D T Q H K L R S A V E E M	72
GAG GTG GAG GAG CTG ATG GAA GAC ACT CAG CAC AAA CTG CGC AGT GCC GTG GAG GAG ATG	325
E A E E A A A K T S S E V N L A S L P P	92
GAG GCG GAA GAA GCA GCT GCT AAA ACG TCC TCT GAG GTG AAC CTG GCA AGC TTA CCT CCC	385
N Y H N E T S T E T R V G N N T V H V H	112
AAC TAT CAC AAT GAG ACC ACC ACG GAG ACC AGG GTG GGA AAT AAC ACA GTC CAT GTG CAC	445
Q E V H K I T N N Q S G Q V V F S E T V	132
CAG GAA GTT CAC AAG ATA ACC AAC AAC CAG AGT GGA CAG GTG GTC TTT TCT GAG ACA GTC	505
I T S V G D E E G K R S H E C I I D E D	152
ATT ACA TCT GTA GGG GAT GAA GAA GGC AAG AGG AGC CAT GAA TGT ATC ATT GAT GAA GAC	565
C G P T R Y C Q F S S F K Y T C Q P C R	172
TGT GGG CCC ACC AGG TAC TGC CAG TTC TCC AGC TTC AAG TAC ACC TGC CAG CCA TGC CGG	625
D Q Q M L C T R D S E C C G D Q L C A W	192
GAC CAG CAG ATG CTA TGC ACC CGA GAC AGT GAG TGC TGT GGA GAC CAG CTG TGT GCC TGG	685
G H C T Q K A T K G G N G T I C D N Q R	212
GGT CAC TGC ACC CAA AAG GCC ACC AAA GGT GGC AAT GGG ACC ATC TGT GAC AAC CAG AGG	745
D C Q P G L C C A F Q R G L L F P V C T	232
GAT TGC CAG CCT GGC CTG TGT GCC TTC CAA AGA GGC CTG CTG TTC CCC GTG TGC ACA	805
P L P V E G E L C H D P T S Q L L D L I	252
CCC CTG CCC GTG GAG GGA GAG CTC TGC CAT GAC CCC ACC AGC CAG CTG CTG GAT CTC ATC	865
T W E L E P E G A L D R C P C A S G L L	272
ACC TGG GAA CTG GAG CCT GAA GGA GCT TTG GAC CGA TGC CCC TGC GCC AGT GGC CTC CTA	925
C Q P H S H S L V Y M C K P A F V G S H	292
TGC CAG CCA CAC AGC CAC AGT CTG GTG TAC ATG TGC AAG CCA GCC TTC GTG GGC AGC CAT	985
D H S E E S Q L P R E A P D E Y E D V G	312
GAC CAC AGT GAG GAG AGC CAG CTG CCC AGG GAG GCC CCG GAT GAG TAC GAA GAT GTT GGC	1045
F I G E V R Q E L E D L E R S L A Q E M	332
TTC ATA GGG GAA GTG CGC CAG GAG CTG GAA GAC CTG GAG CGG AGC CTA GCC CAG GAG ATG	1105

Figure 8B

A	F	E	G	P	A	P	V	E	S	L	G	G	E	E	E	I	*	
GCA	TTT	GAG	GGG	CCT	GCC	CCT	GTG	GAG	TCA	CTA	GGC	GGG	GAG	GAG	GAG	ATT	TAG	350
GCCCGAGACCCAGCTGAGTCACTGGTAGATGTCAATAGAAATGGCTAATTATTTCCCAGGAGTGTCCCCAAGTGTGG																	1159	
AATGGCCGCAGCTCCTCCCAGTAGCTTTCTCTGGCTTGACAAGGTACAGTGCAGTACATTCTTCCAGCCGCCCTG																	1238	
CTTCTCTGACTTGGGAAAGACAGGCATGGCGGTAAAGGCAGCGGTAGTCGTCCTCGCTGTTGCTAGAAACGCTGTC																	1317	
TTGTTCTCATGGATGGAAGATTGTTGAAGGGAGAGGATGGGAAGGGGTGAAGTCTGCTCATGATGGATTGGGGGA																	1396	
TACAGGGAGGAGGATGCCCTGAGACGTGGACTTGGCAAAATGTAACCTTGCTTTGTCTTGCGCCGCTCCCAT																	1475	
GGGCTGAGGCAGTGGCTACACAAGAGCTATGCTGCTCTGTCCTCCACATATTCTACCCCTGTTAGCTCCTACC																	1554	
TCACTGTCAGCACAGCCCTCATAGCCACGCCCCCTCTGCTCACACAGCCTAGGAGGGACCAGAGGGACTTCTCT																	1633	
CAGAGCCCCATGCTCTCTCAACCCATACCAGCCTCTGTGCCAGCGACAGTCCTCCAAATGGAGGGAGTGAAT																	1712	
CCTTTGGTTAATTATTTCTCTCAAGGCACGCCACTAACGGTCAAGGCTAGGCTGACTTGCTATGTCATGTCCTCTAACGTTCG																	1791	
TAGCAGTGTGGTGGACACTGTCTTCCACCGACTGCTCAATACCTCTGAAAGCCAGTGCTGGAGTGCAGTCGTGAA																	1870	
ATTAATTGAGGAAGTATACTTGGCTATTGTAGGGCTAGGATTGTGAATGAAATTGCAAAGTCGCTTAGCAACAAT																	1949	
GGAAAGCCTTCTCAGTCACACCGAGAAGTCACAACCAAGCCAGGTTGTAGAGTACAGCTGTGACATACAGACAGAA																	2028	
GAAGGCTGGCTGGATGTCAGGCCCTCAGATGACGGTTTCAGGTGCCAGGAACATTACCATCTGTATCTATCCAGAGT																	2107	
TATTAATTGAAAGTTGCACACATTGTATAAGCATGCCTTCTCCTGAGTTAAATTATATGTATAACAAACATG																	2186	
TGGCCCTCAAAGATCATGCACAAACCACTACTCTTGCTAATTCTTGGACTTTCTCTTGTATTTCAATAAAATACAAA																	2265	
TCCCCCTCATGCAAAAAAAAAAAAAAGGGCGGCCGC																	2344	
																	2381	